



PCT

RAW SEQUENCE LISTING

DATE: 12/17/2004

PATENT APPLICATION: US/10/516,587

TIME: 14:58:46

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\12172004\J516587.raw

4 <110> APPLICANT: KANEKA CORPORATION
 6 <120> TITLE OF INVENTION: NOVEL ACYLASE GENE
 8 <130> FILE REFERENCE: T719./ACYL-1
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/516,587
 C--> 10 <141> CURRENT FILING DATE: 2004-12-03
 10 <150> PRIOR APPLICATION NUMBER: JP P2002-165722
 11 <151> PRIOR FILING DATE: 2002-06-06
 13 <160> NUMBER OF SEQ ID NOS: 15

ERRORED SEQUENCES

Does Not Comply
 Corrected Diskette Needed
 (pgs 1-2)

470 <210> SEQ ID NO: 15
 471 <211> LENGTH: 26
 472 <212> TYPE: DNA
 473 <213> ORGANISM: Artificial Sequence
 475 <220> FEATURE:
 476 <223> OTHER INFORMATION: Description of Artificial Sequence: MT-218 primer
 478 <400> SEQUENCE: 15
 479 ggaagaagct tgtgcagcac ccggcc 26

E--> 480 1
 E--> 482 1/13
 E--> 484 2/13
 E--> 486 3/13
 E--> 488 4/13
 E--> 490 5/13
 E--> 492 6/13
 E--> 494 7/13
 E--> 496 8/13
 E--> 498 9/13
 E--> 500 10/13
 E--> 502 11/13
 E--> 504 12/13
 E--> 506 13/13

deleted

VERIFICATION SUMMARY

DATE: 12/17/2004

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:480 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=15
L:482 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:484 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:486 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:488 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:490 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:492 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:494 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:496 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:498 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:500 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:502 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:504 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:506 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3
L:506 M:252 E: No. of Seq. differs, <211> LENGTH:Input:26 Found:39 SEQ:15